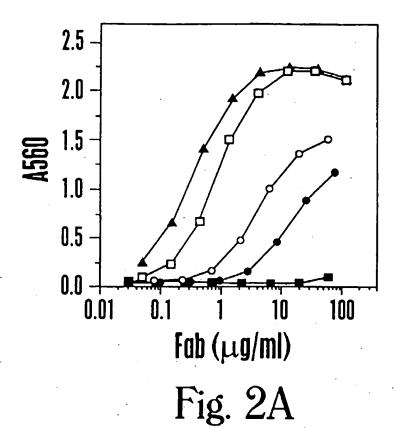
FIGURE 1

VK Dom	ain	
CD40 VKIII	1 10 20 30 40 DIVLTQSPATLSVTPGDRVSLSCRASQSISDYLHWYQQKSHESPRLLIK	
CD40 VKIII/JK4	50 60 70 80 90 YASHSISGIPSRFSGSGSGSDFTLSINSVEPEDVGIYYCQHGHSFPRTFG	
VH Dom CD40 VH7	DIN 1 10 20 30 40 QIQLVQSGPELKKPGETVRISCKASGYAFTTTGMQWVQEMPGKGLKWIG	
CD40 VH7	50 60 70 80 abc 90 WINTHSGVPKYVEDFKGRFAFSLETSANTAYLQISNLKNEDTATYFCVR WINTNTGNPTYAQGFTGRFVFSLDTSVSTAYLQISSLKAEDTAVYYCAR	
CD40 JH4	abcde 110 SGNGNYDLAYFAYWGQGTLVTVSA YFDYWGQGTLVTVSS	



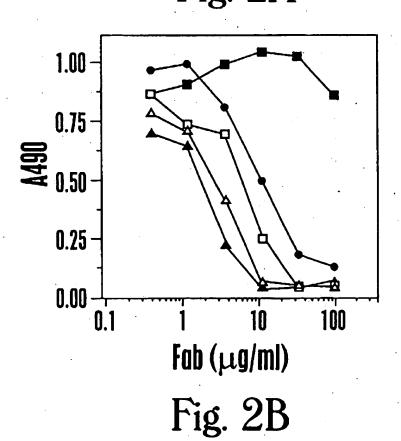
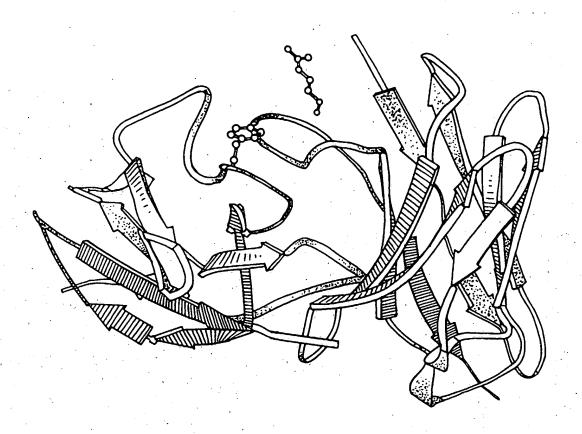


FIGURE 3



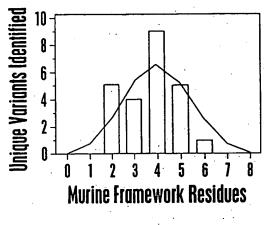


Figure 4A

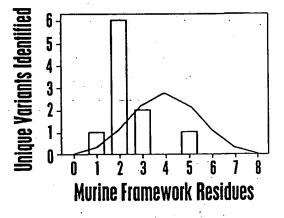


Figure 4B

FIGURE 5

$V_{\rm H} \ {\tt Domain}$

	FR1	CDR-H1	FR2	CDR-H2	FR3
	1 1234567890123456789012345	3 6789012345	4 67890123456789	5 0123456789012345	7 67890123456789012abc345678901234
NMC4	VQLLESGPGLVAPSOSLSITCTVS	GFSLTDYGVD	WVRQPPGKGLEWLG	MIWGDGSTDYNSALKS	RLSITKDNSKSQVFLKMNSLQTDDTARYYCVR
	EVÇLVESGGGLIVQFGGSLKLISCAAS CDR-H3 1 ··· 1		мукфарскопемус		KFTISKUDSKNSLYLQMNSLKTEDIAVYYCAK
	0 567890abcdef12 3456789012		•		
NMC4		WGQGTSVTVS (SEQ ID NO:5)	:5)		
ЈНба	SALALISÕEM	GOGITVIVS (SEQ ID NO:7)	. (2)		

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ł	FR1	CDR-L1	FR2	CDR-1.2	FR3	
	1 12345678901234567890123	3 45678901234	4 567890123456789	5 0123456	1 2 2 3 4 4 5 5 6 7 7 8 12345678901234567890123456789012345678901234567890123456789012345678	78
NMC4 VL	SSLSASLGDRVTISC	SASODINKYLN	WYQQKPDGAVKLLIF	YTSSLHS	SSLSASLGDRVTISC SASQDINKYLN WYQQKPDGAVKLLIF YTSSLHS GVPSRFSGSGSGTDYSLTISNLEPEDIATYYC	χÇ
VKIDPK9/012	VKIDPK9/012 DIQMTQSPS <u>S</u> LSASVGDRVTITC	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	WYQQKPGKAPKLLIY			χÇ
		,		,		

CDR-L3 1 9 0 0 901234567 8901234567 NMC4 VL QQYEKLPWI FGGGTKLEVK (SEQ ID NO:6)

------ FGGGTKVEIK (SEQ ID NO:8).

JK4

vWF Elisa of Combinatorial Positives 0.1 Fab ug/ml 0.01 0.001 0.0001 9.0 0.5 O.D. 560, Dev 30' 0.1 0.2 -0.7

FIGURE 6